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FAX TRANSMISSION COVER SHEET

DATE: 8/23/2006Total Number of Pages Faxed: 9

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NAME:

Becares Rafael D

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PCT

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571-273-0459

FROM: Mark Spencer

Voice Ph. Number: (571)272-2510

FAX Ph. Number: (571)273-0221

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A DOCPHOENIX

STIC INTERNAL

CRFD
Computer Readable Form Defective

CRFE 7
Computer Readable Form "ENTERED"

TC INTERNAL

SEQREQ
Sequence Problem Att. From Exr.

INTERNAL SEQUENCE DOCUMENT INDEX SHEET

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/530,951
Source: IFWP
Date Processed by STIC: 08/25/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 08/25/2006

PATENT APPLICATION: US/10/530,951

TIME: 14:42:24

Input Set : A:\sequence listing ascii.txt

Output Set: N:\CRF4\08252006\J530951.raw

3 <110> APPLICANT: Hariharan, Kandasamy
4 Daniels, Mark
5 McLachlan, Karen
7 <120> TITLE OF INVENTION: GENES OVEREXPRESSED BY OVARIAN CANCER AND THEIR USE IN
DEVELOPING NOVEL
8 THERAPEUTICS
10 <130> FILE REFERENCE: 2159.0620008/EKS/J-H
12 <140> CURRENT APPLICATION NUMBER: 10/530,951
C--> 13 <141> CURRENT FILING DATE: 2005-04-07
15 <150> PRIOR APPLICATION NUMBER: PCT/US03/16253
16 <151> PRIOR FILING DATE: 2003-06-10
18 <150> PRIOR APPLICATION NUMBER: 60/386,748
19 <151> PRIOR FILING DATE: 2002-06-10
21 <150> PRIOR APPLICATION NUMBER: 60/396,141
22 <151> PRIOR FILING DATE: 2002-07-17
24 <150> PRIOR APPLICATION NUMBER: 60/405,319
25 <151> PRIOR FILING DATE: 2002-08-23
27 <150> PRIOR APPLICATION NUMBER: 60/428,274
28 <151> PRIOR FILING DATE: 2002-11-22
30 <160> NUMBER OF SEQ ID NOS: 102
32 <170> SOFTWARE: PatentIn version 3.3
34 <210> SEQ ID NO: 1
35 <211> LENGTH: 473
36 <212> TYPE: DNA
37 <213> ORGANISM: Homo sapiens
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42 ccgaataactg ggtaatecca agtttttact attcatgtaa tttttaaaaa taggtaaaag 120
44 ctagtttttat accagccctc gaaagttgta gagcagctgt atctggtagg aagaatccgt 180
46 actcaatgaa aatgtcgggtg cagacagaac aggccttcagt agatgtttat agatgttttc 240
48 acatgtttaca aattaaagta tattcatctc cagtgcacaa agactaattc tgtattgaag 300
50 agtccctcag ctgctagggt gaccagcagc ccaggcctga gggctttcct ggaacctgaa 360
52 acttttcaatg ctaaaactag gacagtccca ggccaaccag gatgggtggg cattcttcta 420
54 gttgctgaga ttttgaaatc aatatttagt gttcattcta ctgttattgg aat 473
57 <210> SEQ ID NO: 2
58 <211> LENGTH: 2992
59 <212> TYPE: DNA
60 <213> ORGANISM: Homo sapiens
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64 <221> NAME/KEY: misc_feature
65 <223> OTHER INFORMATION: n is a, c, g, or t
67 <220> FEATURE:
68 <221> NAME/KEY: misc_feature
69 <222> LOCATION: (637)..(637)

CP8-67

RAW SEQUENCE LISTING

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72 <220> FEATURE:
73 <221> NAME/KEY: misc_feature
74 <222> LOCATION: (1192)..(1192)
75 <223> OTHER INFORMATION: n is a, c, g, or t
77 <220> FEATURE:
78 <221> NAME/KEY: misc_feature
79 <222> LOCATION: (1399)..(1399)
80 <223> OTHER INFORMATION: n is a, c, g, or t
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83 <221> NAME/KEY: misc_feature
84 <222> LOCATION: (1436)..(1437)
85 <223> OTHER INFORMATION: n is a, c, g, or t
87 <220> FEATURE:
88 <221> NAME/KEY: misc_feature
89 <222> LOCATION: (1447)..(1447)
90 <223> OTHER INFORMATION: n is a, c, g, or t
92 <220> FEATURE:
93 <221> NAME/KEY: misc_feature
94 <222> LOCATION: (1453)..(1453)
95 <223> OTHER INFORMATION: n is a, c, g, or t
97 <220> FEATURE:
98 <221> NAME/KEY: misc_feature
99 <222> LOCATION: (1460)..(1460)
100 <223> OTHER INFORMATION: n is a, c, g, or t
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105 <223> OTHER INFORMATION: n is a, c, g, or t
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108 <221> NAME/KEY: misc_feature
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114 <222> LOCATION: (1529)..(1531)
115 <223> OTHER INFORMATION: n is a, c, g, or t
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129 <222> LOCATION: (1549)..(1549)
130 <223> OTHER INFORMATION: n is a, c, g, or t

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Input Set : A:\sequence listing ascii.txt

Output Set: N:\CRF4\08252006\J530951.raw

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139 <222> LOCATION: (1564)..(1564)
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147 <220> FEATURE:
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149 <222> LOCATION: (1574)..(1575)
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154 <222> LOCATION: (1577)..(1577)
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158 <221> NAME/KEY: misc_feature
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160 <223> OTHER INFORMATION: n is a, c, g, or t
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164 <222> LOCATION: (1588)..(1588)
165 <223> OTHER INFORMATION: n is a, c, g, or t
167 <220> FEATURE:
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169 <222> LOCATION: (1598)..(1598)
170 <223> OTHER INFORMATION: n is a, c, g, or t
172 <220> FEATURE:
173 <221> NAME/KEY: misc_feature
174 <222> LOCATION: (1600)..(1601)
175 <223> OTHER INFORMATION: n is a, c, g, or t
177 <220> FEATURE:
178 <221> NAME/KEY: misc_feature
179 <222> LOCATION: (1604)..(1604)
180 <223> OTHER INFORMATION: n is a, c, g, or t
182 <220> FEATURE:
183 <221> NAME/KEY: misc_feature
184 <222> LOCATION: (1606)..(1606)
185 <223> OTHER INFORMATION: n is a, c, g, or t
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188 <221> NAME/KEY: misc_feature
189 <222> LOCATION: (1608)..(1608)
190 <223> OTHER INFORMATION: n is a, c, g, or t
192 <220> FEATURE:

RAW SEQUENCE LISTING

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Input Set : A:\sequence listing ascii.txt

Output Set: N:\CRF4\08252006\J530951.raw

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193 <221> NAME/KEY: misc_feature
194 <222> LOCATION: (1612)..(1612)
195 <223> OTHER INFORMATION: n is a, c, g, or t
197 <220> FEATURE:
198 <221> NAME/KEY: misc_feature
199 <222> LOCATION: (1623)..(1623)
200 <223> OTHER INFORMATION: n is a, c, g, or t
202 <220> FEATURE:
203 <221> NAME/KEY: misc_feature
204 <222> LOCATION: (1625)..(1625)
205 <223> OTHER INFORMATION: n is a, c, g, or t
207 <220> FEATURE:
208 <221> NAME/KEY: misc_feature
209 <222> LOCATION: (1640)..(1641)
210 <223> OTHER INFORMATION: n is a, c, g, or t
212 <220> FEATURE:
213 <221> NAME/KEY: misc_feature
214 <222> LOCATION: (1649)..(1649)
215 <223> OTHER INFORMATION: n is a, c, g, or t
217 <400> SEQUENCE: 2
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220 gccagtcagt ggttggtggt cagttatcag gacaggacgg aattctttgt caggccgctc      120
222 agtttttcag gtgaacctgt gataagggag tggagtctgg ccacgggtgc aggtgatctg      180
224 ctgaagtcca tggaaggatc ttccagtcct tgtttttcag atattgggtt ctgtttatct      240
226 cttagaaagt ctggttaata cacaaagagg gggtagctag gcatactctaa cctcctatct      300
228 catcaaagcc agggaaactt aaagtttttc tgggtgatct gggccaagag ggggtccattc      360
230 agtctgtcag aggcattcaga ttctattctg gttccacagg ggactccaga acgtgattcc      420
232 cggctctctc accttaatcc tgtcatcacc ctaggagatt ttagtggcaa ttagatagat      480
234 tcttcctagc tctgagctca aagtttccca aagggtattt ttctagtatt gagagacatt      540
236 aatgggtgta gggatagaga gattaaatga gacctatgat cacataagtt tggaaaatga      600
W--> 238 tgaataagca aagggtaatg tagaagaaga aatttttnatt taaaaccaaa gctgatgatg      660
240 tggattccta ataagataag tgagcaacaa cttggagggg gcccaggtg gggagagcaa      720
242 tttttctgcc agacagctca tcactgacaa cccactggca ggacatcctg ttcccaata      780
244 cctcactcca cacatagccc caacagcaca acctcattct gcacataccc cctccagtac      840
246 aacctagaa aacttccttc cagtcctctg ctctttgncg acagcccctt ctctgccgtg      900
248 ctgcccattg cctccttgca acatattctc cctgtaatga gcttgctttc ttaactcac      960
250 tactgtcttg gtaaatctt ttaccatctg tgacgccagc cccagccagt tgcacctgca     1020
252 acagatgaca gctacttctg tcttctaagg aggggtggaat ggttcacatc gataagaaaa     1080
254 tacgtactcc atggctccat tccacttacg gggactggag tggggaggag aaagtagaaa     1140
W--> 256 ggaaactaac atttctgtgc ttctgctgt tttctatggt gatgacctgc angtgcaggg     1200
258 cagacagccc atgtgacctg attgcctaag attttctctg tgtacacctg ttggccagca     1260
260 aagttattac taggccact ttcaatctta aaagtgtcat agtttgagc acgtattatg     1320
262 tggttacctg ttaatgaggc ttgaaaaatg taaaagtagg ttcaagacca aataaggaga     1380
W--> 264 ggaaagcaaa gttgttttng tacaataaag tgaggagagt tgagttgggg tagggungga     1440
W--> 266 gggggcnagg ganataaaan gaaagttttg nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn     1500
W--> 268 nnnnnnnnnn nnnnnnnnnn chgcctaann ncanaataaa gggggnaant ggtaaaaagg     1560
W--> 270 tncnctnnaa aatnngntgg anaatttnaa agggaggntn naantnanag tngaaagtga     1620
W--> 272 cntnaaaaaa aataaacccn nctggaaant tctcagaatc tcaggatggg cccagagta     1680
274 tctaaagatg ctacagttca agggattgag ccaattgtat ataaatctta atggataggt     1740
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276 tgacctcagc ataaaacttg ggtggaaatt ttaaacaggt ttctttatatt cagcactttct 1800
278 cagagcaactc attgtataag gtactttgtg aatatccaga tagtattctt caaactctct 1860
280 tttattttccc caggggggcat cccataggac aagaagcatt ctttgtgaca ctctgtggga 1920
282 agagctgggt taaaggggta cctgtctggg caacactgtc ccacaggggc ccccatgacc 1980
284 aaactaactc tgctttctacc cagaaagggt gcagagtagc cactagactt ttatgtggca 2040
286 aatgggatgg ttatgcccag cctgaagcca agatgccctt tctggttgcc ttgattttgtg 2100
288 tttaacagct ccaaagtctt aatgaggcag taagagacgt ctctcttggg cagtacttcc 2160
290 caactagggg tgagtttgcc acccttacct ccatcccagt gaatatttgc aattcctaaa 2220
292 gacgtgtttt gattgtcaca cctgggtggg gaacatgcta ctggcatcta atgcatagag 2280
294 ggcagtaatg ctgctaaaca tctttcaacg cacagggaca gagccccaca aaaaagaatt 2340
296 atctaggccc caaatgtcca taacactgct gttgagaaaa acctaccgca ggatcttact 2400
298 gggcttcata ggtaagottg cctttgttct ggcttctgta gatataataa ataaagacac 2460
300 tgcccagtc cctccctcaac gtcccagagc agggctcaag gcaattccaa taacagtaga 2520
302 atgaacacta aatatttgatt tcaaaatctc agcaactaga agaatgacca accatcctgg 2580
304 ttggcctggg actgtcctag ttttagcatt gaaagtttca ggttccagga aagccctcag 2640
306 gcttgggctg ctgggtcacc tagcagctga gggactcttc aatacagaat tagtctttgt 2700
308 gcactggaga tgaatatact ttaattttgta acatgtgaaa acatctataa acatctactg 2760
310 aagcctgttc tgtctgcacc gacattttca ttgagtacgg attcttctta ccagatacag 2820
312 ctgctctaca actttcgagg gctgggtataa aactagcttt tacctatttt taaaaattac 2880
314 atgaatagta aaaacttggg ttaaccaggt attcgggtat ttcaatctt ccttgggagc 2940
316 tttagaggag gacaaaataa aagattattt caacatcaaa aaaaaaaaaa aa 2992
319 <210> SEQ ID NO: 3
320 <211> LENGTH: 477
321 <212> TYPE: DNA
322 <213> ORGANISM: Homo sapiens
324 <400> SEQUENCE: 3
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327 aaaccagcag atgottgccca cttcccaagg tgtaaaaatg tgaagattaa ggtaaactga 120
329 atgtcgagga gtgtaaagag atggcaaaac acagataaaa acatccaaaa agcctctggg 180
331 ggcagggtcaa gcttatgatt caacagttag aaaacaaaaa ttacttggac atccccttct 240
333 acttaaagtg atatactgga attgaaaata ttaactgtta gttttagaaa ctaagattac 300
335 ttgaagtagg ctcatccag aatgcttttc ttttttcctt cctgaaacaa ttacatcaaa 360
337 cttagatata ctaatgttta ttttagata tactccttaa aggcatttat gttcacccct 420
339 ttcgaggatg agaaatttac attaactaat aaacttaaca ttgatcttaa gaactgg 477
342 <210> SEQ ID NO: 4
343 <211> LENGTH: 340
344 <212> TYPE: DNA
345 <213> ORGANISM: Homo sapiens
347 <400> SEQUENCE: 4
348 tttttttttt ttttttttcc aggattccaa attttatttt ttaaaaaatt gaaaaaacac 60
350 acccaggaca acattttcttt gatcaataaa ctttcaggaa atggaggaag ctgttttggg 120
352 acacattcaa agctagttaa cttgaacttg gaaatagggt ttgacaatc caactatggg 180
354 aaacaaatct ctgaacaaat tttaaatgaa acctcacccc cccaaactgt tcaagtggca 240
356 gacaaaaata attaccataa attatatgcc aacacacctt ttaaaaaaca acaacagcaa 300
358 caacaaaaac ccaggagtct gaggatttcc ttagctcctc 340
361 <210> SEQ ID NO: 5
362 <211> LENGTH: 1121
363 <212> TYPE: DNA
364 <213> ORGANISM: Homo sapiens
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/530,951

DATE: 08/25/2006
TIME: 14:42:25

Input Set : A:\sequence listing ascii.txt
Output Set: N:\CRF4\08252006\J530951.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 637,1192,1399,1436,1437,1447,1453,1460,1471,1472,1473,1474
Seq#:2; N Pos. 1475,1476,1477,1478,1479,1480,1481,1482,1483,1484,1485,1486
Seq#:2; N Pos. 1487,1488,1489,1490,1491,1492,1493,1494,1495,1496,1497,1498
Seq#:2; N Pos. 1499,1500,1501,1502,1503,1504,1505,1506,1507,1508,1509,1510
Seq#:2; N Pos. 1511,1512,1513,1514,1515,1516,1517,1518,1519,1520,1522,1529
Seq#:2; N Pos. 1530,1531,1534,1546,1549,1562,1564,1567,1568,1574,1575,1577
Seq#:2; N Pos. 1582,1588,1598,1600,1601,1604,1606,1608,1612,1623,1629,1640
Seq#:2; N Pos. 1641,1649
Seq#:8; N Pos. 319
Seq#:12; N Pos. 434
Seq#:13; N Pos. 238
Seq#:17; N Pos. 444,447,499
Seq#:24; N Pos. 233
Seq#:25; N Pos. 387,395,398,439,460,463

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:35,36,47,48,49,50,51,52

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/530,951

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Input Set : A:\sequence listing ascii.txt

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:238 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:600
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1140
L:264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1380
L:266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1440
L:268 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1500
L:270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1560
L:272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1620
L:510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:300
L:712 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:420
L:735 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:180
L:897 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:420
L:899 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:480
L:1376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:180
L:1411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:360
L:1413 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:420